

OM protein - protein search, using sw model

Run on: April 14, 2006, 20:39:58 ; Search time 190 Seconds
(without alignments)
1244.136 Million cell updates/sec

Title: US-10-796-669A-3

Perfect score: 2769

Sequence: 1 MTKKPARKILSFSTTMRNPK.....FIKKVENSIEWLQINDHLKA 538

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 2443163 seqs, 439378781 residues

Total number of hits satisfying chosen parameters: 2443163

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : A_Geneseq_21:*
1: geneseqp1980s:*

2: geneseqp1990s:*

3: geneseqp2000s:*

4: geneseqp2001s:*

5: geneseqp2002s:*

6: geneseqp2003as:*

7: geneseqp2003bs:*

8: geneseqp2004s:*

9: geneseqp2005s:*

GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: April 14, 2006, 20:43:48 ; Search time 43 Seconds
(without alignments)
1203.828 Million cell updates/sec

Title: US-10-796-669A-3

Perfect score: 2769

Sequence: 1 MTKKPARKILSFSTTMRNPK.....FIKKVENSIEWLQINDHLKA 538

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 283416 seqs, 96216763 residues

Total number of hits satisfying chosen parameters: 283416

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : PIR_80:*

1: pirl:*

2: pir2:*

3: pir3:*

4: pir4:*

GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: April 14, 2006, 20:40:28 ; Search time 235 Seconds
(without alignments)
1615.209 Million cell updates/sec

Title: US-10-796-669A-3

Perfect score: 2769

Sequence: 1 MTKKPARKILSFSTTMRNPK.....FIKKVENSIEWLQINDHLKA 538

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 2166443 seqs, 705528306 residues

Total number of hits satisfying chosen parameters: 2166443

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : UniProt_05.80:*

1: uniprot_sprot:*

2: uniprot_trembl:*